

REPLACEMENT PARAGRAPH 0021

[0021] proteolytically active fragment of the light chain of a Clostridium botulinum toxin, in particular a toxin of the type A, B, C1, D, E, F, and G, characterized by containing the sequence of SEQ ID NO:1 His-Xaa-Xaa-Xaa-His-Xaa-Xaa-His, where Xaa can be any amino acid (for example, Xaa can be Leu, Ile, Val, Arg, Met);

- 2 -

3/5/04: Amd for Ser. No. 10/084,903 - Inventor(s): Bigalke et al. - Filing Date: 8/27/2002

REPLACEMENT PARAGRAPH 0023

[0023] proteolytically active fragment of the light chain of the tetanus toxin, characterized by containing the sequence of SEQ ID NO:2 His-Asp-Leu-Ile-His-Val-Leu-His;

- 3 -

3/5/04: Amd for Ser. No. 10/064,803 - Inventor(s): Bigalke et al. - Filing Date: 8/27/2002

REPLACEMENT PARAGRAPH 0026

[0026] In the sequence of SEQ ID NO:1 His-Xaa-Xaa-Xaa-His-Xaa-Xaa-His of the proteolytically active fragment of the light chain of a Clostridium botulinum toxin, the first Xaa is preferably Asp or Ala; the second Xaa is preferably Leu or Ile; the third Xaa is preferably Ile, Asn or Tyr; and the fourth and fifth Xaa are preferably an amino acid with a non-polar functional group, i.e., amino acids such as Val, Leu, Ile, or Ala, respectively.